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03-01 0280

OIPE

JF R

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/817,360

DATE: 04/05/2001
TIME: 12:14:04

Input Set : A:\SeqList for UCSF-129CIP.txt
Output Set: N:\CRF3\04052001\I817360.raw

ENTERED

4 <110> APPLICANT: German, Michael S.
5 Lin, Joseph
7 <120> TITLE OF INVENTION: PRODUCTION OF PANCREATIC ISLET CELLS
8 AND DELIVERY OF INSULIN
13 <130> FILE REFERENCE: UCSF-129CIP
C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/817,360
C--> 15 <141> CURRENT FILING DATE: 2001-03-20
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16 <151> PRIOR FILING DATE: 2000-03-24
18 <150> PRIOR APPLICATION NUMBER: 60/128,180
19 <151> PRIOR FILING DATE: 1999-04-06
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33 tctaacttgc agttaaataaa atcaggcaag gctggctat gaggcagaca agtgtgaaga 180
34 aggagaagga ggaggagaag gagaaggaga aagaagaaga aggaggagaa gaagaagaag 240
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37 gacccccaga ctcaggactt cctctatttt ctgcattttt gggctctttg tttgccttg 420
38 aaaaaaaaaatg ttttctccca aatcaaggag cagtagctgg tgcaaggaa aatctagggc 480
39 taggagtctt aagatatgac ttctatgtgg ttctgataga acttgctggg tgacccttgag 540
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41 cattctttag agaagaagac aagctcttag tgagtggta cctaaggagc cagtcgcgc 660
42 aaaatctaa cctgacagtc ccagatggc ccttatttg ttctgaccct ggtctcaggc 720
43 ttcatttccc cacagcaagg gaaggagcct gctcacagag caccagctaa gatcagcagg 780
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46 gggcagactc acagacccatc ctcatcccc tacagctctg aagtccggc 960
47 ctgtccccctc ctgcagttt cgggagactc agatatctg gacgtctgt aaagagaagc 1020
48 cttccctcgcc taaggagact taaacccggga tacttaaacc tcccgctcg gcgtttccct 1080
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50 ggaagcagaa gaccacggg tgcctccaggc ggggacaaga ggagggctg gggaaagaaag 1200
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58 gtctcagcta ctgggaaggc tgaagtgggaa ggatagctt agccctggag atcaaggctg 1680
59 cagtgagctg tgattgcacc actgcagttcc agcctggcg acagaaggag accgttttt 1740

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62	gggatactat	ctacaagctg	tagtaggctt	gtagtaatgg	aatgtccgct	tgaggggtcc	1920
63	ccgcacagcc	aaccccccggcc	tctggagtgg	gatctatggg	ggtgggggtc	taagcgctc	1980
64	tggggagtgt	gaggtagcat	ctcagggtgt	ggcagaggct	cggacacccc	caaaaggct	2040
65	gtgaatggaa	gggacatagg	caggatctct	ctcagtgtat	tcccctgtct	tccaggatga	2100
66	agagagggcag	tgaaacacca	ggagacagg	gctgtcttta	gaattcctgg	acccttctcc	2160
67	aggctgttag	tcaggacaat	gagctgtgg	ttgtctttgc	cactatcttc	ctgtgcgatt	2220
68	tcagacaagg	caccccccctc	actaagccta	aatttcccca	tgtgtacgt	gcagggattt	2280
69	taccctagag	gcatcaaagt	cccctccagg	acagatgcta	aggaaagata	ggcttaggagc	2340
70	aaagccgtct	gagggtggct	gaccagagcc	acacgaggct	cttctcaact	ggcgaggctc	2400
71	tttggaggaac	cgagagttgc	tgggaccctag	cccgccctcg	agagagcaaa	cagagcggcg	2460
72	ctcccccctcc	ccgaccccccgg	cccttgc	ggaatccagc	tgtgtcg	gggaggagcg	2520
73	ggctcgctg	gcccggcccc	agggcccccgg	cgctgattgg	ccgggtggc	gggcagcagc	2580
74	ccggcaggca	cgctccctggc	ccgggcaag	cagataaagc	gtgccaaggg	gcacacgact	2640
75	tgtgtctcg	gaaatccctg	cggtctcacc	ggcgcgcctc	gagagagagc	gtgacagagg	2700
76	cctcggaccc	cattctctct	tcttttctcc	tttggggctg	gggcaactcc	caggcgggggg	2760
77	cgcctgcagc	ttagctgaac	ttggcgacca	gaagcccgct	gagctccca	cggccctcg	2820
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86	aatcgaatgc	acaacctcaa	ctcggcaact	gacgcctgc	gcccgtt	gcccac	3360
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104	ttttaaatct	gtttgttaat	tatatgtgc	ctttgttgt	caattttgt	acagtaaaat	4440
105	tatatggccc	ctccccctgtt	ttacacattt	gtattttata	atgagattt	acagcaggga	4500
106	aaagcctata	ttttggat	tagattttt	aggattgt	ggatgacatt	taagccaata	4560
107	aaaaaaaatg	gaccccaag	aagccctggc	aagatgactc	cattgtgt	tggggagagg	4620
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117 aaaacttatt aagaactgt tcaaggttt cagccacacc atgtcttta ctggcaaggt 5220
118 ggaataggac tggtgcagca tgagcaactga aatctgtccc aggagtgcga gttagagcacc 5280
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128 Thr Glu Arg Ser Phe Pro Arg Ala Ser Glu Asp Glu Val Thr Cys Pro
129 20 25 30
130 Thr Ser Ala Pro Pro Ser Pro Thr Arg Thr Arg Gly Asn Cys Ala Glu
131 35 40 45
132 Ala Glu Glu Gly Gly Cys Arg Gly Ala Pro Arg Lys Leu Arg Ala Arg
133 50 55 60
134 Arg Gly Gly Arg Ser Arg Pro Lys Ser Glu Leu Ala Leu Ser Lys Gln
135 65 70 75 80
136 Arg Arg Ser Arg Arg Lys Lys Ala Asn Asp Arg Glu Arg Asn Arg Met
137 85 90 95
138 His Asn Leu Asn Ser Ala Leu Asp Ala Leu Arg Gly Val Leu Pro Thr
139 100 105 110
140 Phe Pro Asp Asp Ala Lys Leu Thr Lys Ile Glu Thr Leu Arg Phe Ala
141 115 120 125
142 His Asn Tyr Ile Trp Ala Leu Thr Gln Thr Leu Arg Ile Ala Asp His
143 130 135 140
144 Ser Leu Tyr Ala Leu Glu Pro Pro Ala Pro His Cys Gly Glu Leu Gly
145 145 150 155 160
146 Ser Pro Gly Gly Ser Pro Gly Asp Trp Gly Ser Leu Tyr Ser Pro Val
147 165 170 175
148 Ser Gln Ala Gly Ser Leu Ser Pro Ala Ala Ser Leu Glu Glu Arg Pro
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157 <212> TYPE: DNA
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166	gaggccttcc	ttatatatat	ataggcaccc	ccaaacctcc	ttcatgctac	caagaaaaggg	360										
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216 His Asn Tyr Ile Trp Ala Leu Thr Gln Thr Leu Arg Ile Ala Asp His
217 130 135 140
218 Ser Phe Tyr Gly Pro Glu Pro Pro Val Pro Cys Gly Glu Leu Gly Ser
219 145 150 155 160
220 Pro Gly Gly Ser Asn Gly Asp Trp Gly Ser Ile Tyr Ser Pro Val
221 165 170 175
222 Ser Gln Ala Gly Asn Leu Ser Pro Thr Ala Ser Leu Glu Glu Phe Pro
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242 <212> TYPE: DNA
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248 <400> SEQUENCE: 6
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L: 15 M: 270 C: Current Application Number differs, Replaced Current Application No
L: 15 M: 271 C: Current Filing Date differs, Replaced Current Filing Date